

MILNVLRICI IVCLVNDGAG KHSEGRERTK TYSLNSRGYF 40
RKERGARRSK ILLVNTKGLD EPHIGHGDFG LVAELFDSTR 80
THTNRKEPDM NKVKLFSTVA HGNKSARRKA YNGSRRNIFS 120
RRSFDKRNTE VTEKPGAKMF WNNFLVKMNG APQNTSHGSK 160
AQEIMKEACK TLPFTQNI VH ENCDRMVIQN NLCFGKCISL 200
HVPNQQDRRN TCSHCLPSKF TLNHLTLNCT GSKNVVKVVM 240
MVEECTCEAH KSNFHQTAQF NMDTSTTLHH 270

Figure 1. Deduced amino acid sequence of *Xenopus cerberus* protein. SEQ ID NO:1.

Figure 2. Nucleotide sequence of the full-length cerberus DNA derived from the *Xenopus* organizer. The sense strand is on top (in the 5' to 3' direction) and the antisense strand on the bottom line (on the opposite direction). SEQ ID NO:2.

GAATCCAG CAAGTCGCTC AGAAACACTG CAGGGTCTAG ATATCATACA ATGTTACTAA	60
CTTAAGGGTC GTTCAGCGAG TCTTGTGAC GTCCAGATC TATAGTATGT TACAATGATT	
ATGTA CTAGACATAA TAGCAGACGG AACACTTACT ACCTCGTCTT TTTGTGAGTC	120
AAGGACGAGA AAGGACAAAA ACATATTACAC TAAACAGCAG AGGTTACTTC AGAAAAGAAA	180
TTCCCTGCTCT TTCCCTGTTT TGTATAAGTG AATTGTCGTC TCCAATGAAG TCTTTCTTT	
GAGGAGCACG TAGGAGCAAG ATTCTGCTGG TGAATACTAA AGGTCTTGAT GAACCCCA	240
CTCCTCGTGC ATCCTCGTTC TAAGACGACC ACTTATGATT TCCAGAACTA CTTGGGGTGT	
TTGGGCATGG TGATTTCCG TTAGTAGCTG AACTATTGTA TTCCACCAGA ACACATACAA	300
AACCCGTACCC ACTAAAAGCG AATCATCGAC TTGATAAACT AAGGTGGTCT TGTGTATGTT	
ACAGAAAAGA GCCAGACATG AACAAAGTCA AGCTTTCTC AACAGTTGCC CATGGAAACA	360
TGTCTTTCT CGGTCTGTAC TTGTTTCAGT TCGAAAAGAG TTGTCAACGG GTACCTTTGT	
AAAGTGCAAG AAGAAAAGCT TACAATGGTT CTAGAAGGAA TATTTTCCT CGCCGTTCTT	420
TTTCACGTTTC TTCTTTCGA ATGTTACCAA GATCTTCCTT ATAAAAAGGA GCGGCAAGAA	
TTGATAAAAG AAATACAGAG GTTACTGAAA AGCCTGGTGC CAAGATGTT TGGAAACAATT	480
AACTATTTTC TTTATGTCTC CAATGACTTT TCGGACCACG GTTCTACAAG ACCTTGTAA	
TTTTGGTTAA AATGAATGGA GCCCCACAGA ATACAAGCCA TGGCAGTAAA GCACAGGAAA	540
AAAACCAATT TTACTTACCT CGGGGTGTCT TATGTTCGGT ACCGTCATTT CGTGTCCCTT	
TAATGAAAGA AGCTTGCAAA ACCTTGTCTT TCACTCAGAA TATTGTACAT GAAAACGTG	600
ATTACTTTCT TCGAACGTTT TGGAAACAAAA AGTGAGTCTT ATAACATGTA CTTTGACAC	
ACAGGATGGT GATACAGAAC AATCTGCTG TTGGTAAATG CATCTCTCTC CATGTTCCAA	660
TGTCCTACCA CTATGTCTG TTAGACACGA AACCATTTAC GTAGAGAGAG GTACAAGGTT	
ATCAGCAAGA TCGACGAAAT ACCTGTTCCC ATTGCTTGCC GTCCAAATTT ACCCTGAACC	720
TAGTCGTTCT AGCTGCTTTA TGAACAAGGG TAACGAACGG CAGGTTAAA TGGGACTTGG	
ACCTGACGCT GAATTGTACT GGATCTAAGA ATGTTAGTAAA GGTTGTCTG ATGGTAGAGG	780
TGGACTGCGA CTTAACATGA CCTAGATTCT TACATCATT CCAACAGTAC TACCATCTCC	
AATGCACGTG TGAAGCTCAT AAGAGCAACT TCCACCAAAAC TGCACAGTTT AACATGGATA	840
TTACGTGCAAC TTCTCGAGTA TTCTCGTTGA AGGTGGTTG ACGTGTAAA TTGTACCTAT	
CATCTACTAC CCTGCACCAT TAAAGGACTG CCATACAGTA TGGAAATGCC CTTTTGTTGG	900
GTAGATGATG GGACGTGGTA ATTTCCGTGAC GGTTATGTCT ACCTTTACGG GAAAACAACC	
AATATTGTT ACATACTATG CATCTAAAGC ATTATGTTGC CTTCTATTTC ATATAACCAC	960
TTATAAACAA TGTATGATAC GTAGATTCG TAATACAACG GAAGATAAAAG TATATTGGTG	
ATGGAATAAG GATTGTATGA ATTATAATTA ACAAAATGGCA TTTTGTGTAACATGCAAGAT	1020
TACCTTATTC CTAACATACT TAATATTAAT TGTTTACCGT AAAACACATT GTACGTTCTA	

CTCTGTTCCA TCAGTTGCAA GATAAAAGGC AATATTTGTT TGACTTTTT TCTACAAAAT GAGACAAGGT AGTCAACGTT CTATTTCCG TTATAAACAA ACTGAAAAAA AGATGTTTA	1080
GAATACCCAA ATATATGATA AGATAATGGG GTCAAAACTG TTAAGGGTA ATGTAATAAT CTTATGGGTT TATATACTAT TCTATTACCC CAGTTTGAC AATTCCCCAT TACATTATTA	1140
AGGGACTAAG TTTGCCAGG AGCAGTGACC CATAACAACC AATCAGCAGG TATGATTTAC TCCCTGATTC AAACGGGTCC TCGTCACTGG GTATTGTTGG TTAGTCGTCC ATACTAAATG	1200
TGGTCACCTG TTTAAAAGCA AACATCTTAT TGTTGCTAT GGGTTACTGC TTCTGGCAA ACCAGTGGAC AAATTTCGT TTGTAGAATA ACCAACGATA CCCAATGACG AAGACCCGTT	1260
AATGTGTGCC TCATAGGGGG GTTAGTGTGT TGTGTACTGA ATAAATTGTA TTTATTCAT TTACACACGG AGTATCCCC CAATCACACA ACACATGACT TATTTAACAT AAATAAAGTA	1320
TGTTACAAA AAAAAAAA ACAATGTTT TTTTTTTT	

Fig. 2. (Continuation page 2, SEQ ID NO:2).

MSRTRKVDSL LLLAIPGLAL LLLPNAYCAS CEPVRIPMCK SMPWNMTKMP NHLHHSTQAN 60
AILAIEQFEG LLTTECSQDL LFFLCAMYAP ICTIDFQHEP IKPCKSVcer ARAGCEPILI 120
KYRHTWPESL ACEELPVYDR GVCISPEAIV TVEQGTDsMP DFSMDSNNGN CGSGREHCKC 180
KPMKATQKTY LKNNNYNYVIR AKVKEVKVKC HDATAIVEVK EILKSSLVNI PKDTVTLYTN 240
SGCLCPQLVA NEEYIIMGYE DKERTRLLL V EGSLAEKWRD RLAKKVKRWD QKLRRPRKSK 300
DPVAPIPNKN SNSRQARS

Figure 3. Deducing amino acid sequence of Xenopus frazzled protein. SEQ ID NO:3.

Figure 4. Nucleotide sequence of the full-length frazzled cDNA derived from the Xenopus organizer. The sense strand of the DNA on top (5' to 3' direction) and the antisense strand on the bottom line (opposite direction). SEQ ID NO:4.

GAATTCCCTT TCACACAGGA CTCCCTGGCAG AGGTGAATGG TTAGCCCTAT GGATTTGGTT	60
CTTAAGGGAA AGTGTGTCTT GAGGACCGTC TCCACTTACC AATCGGGATA CCTAAACCAA	
TGTTGATTT GACACATGAT TGATTGCTTT CAGATAGGAT TGAAGGACTT GGATTTTTAT	120
ACAACTAAAA CTGTGTACTA ACTAACGAAA GTCTATCCTA ACTTCCTGAA CCTAAAATA	
CTAATTCTGC ACTTTTAAAT TATCTGAGTA ATTGTTCATT TTGTATTGGA TGGGACTAAA	180
GATTAAGACG TGAAAATTAA ATAGACTCAT TAACAAGTAA AACATAACCT ACCCTGATTT	
GATAAACTTA ACTCCTTGCT TTTGACTTGC CCATAAACTA TAAGGTGGGG TGAGTTGTAG	240
CTATTGAAT TGAGGAACGA AAACTGAACG GGTATTTGAT ATTCCACCCC ACTCAACATC	
TTGCTTTAC ATGTGCCAG ATTTCCCTG TATTCCCTGT ATTCCCTCTA AAGTAAGCCT	300
AACGAAAATG TACACGGGTC TAAAAGGGAC ATAAGGGACA TAAGGGAGAT TTCATTGGA	
ACACATACAG GTTGGGCAGA ATAACAATGT CTCGAACAAG GAAAGTGGAC TCATTACTGC	360
TGTGTATGTC CAACCCGTCT TATTGTTACA GAGCTTGTTC CTTTCACCTG AGTAATGACG	
TACTGGCCAT ACCTGGACTG GCGCTTCTCT TATTACCCAA TGCTTACTGT GCTTCGTGTG	420
ATGACCGGTA TGGACCTGAC CGCGAAGAGA ATAATGGGTT ACGAATGACA CGAAGCACAC	
AGCCTGTGCG GATCCCCATG TGCAAATCTA TGCCATGGAA CATGACCAAG ATGCCCAACC	480
TCGGACACGC CTAGGGTAC ACGTTTAGAT ACGGTACCTT GTACTGGTTC TACGGGTTGG	
ATCTCCACCA CAGCACTCAA GCCAATGCCA TCCTGGCAAT TGAACAGTTT GAAGGTTTGC	540
TAGAGGTGGT GTCGTGAGTT CGGTTACGGT AGGACCGTTA ACTTGTCAAA CTTCCAAACG	
TGACCACTGA ATGTAGCCAG GACCTTTGT TCTTCTGTG TGCCATGTAT GCCCCCCATT	600
ACTGGTGACT TACATCGGTC CTGGAAAACA AGAAAGACAC ACGGTACATA CGGGGGTAAA	
GTACCATCGA TTTCCAGCAT GAACCAATTAA AGCCTTGCAA GTCCGTGTGC GAAAGGGCCA	660
CATGGTAGCT AAAGGTGCTA CTTGTTAAT TCGGAACGTT CAGGCACACG CTTTCCCGGT	
GGGCCGGCTG TGAGCCCATT CTCATAAAAGT ACCGGCACAC TTGGCCAGAG AGCCTGGCAT	720
CCCGGCGGAC ACTCGGTAA GAGTATTCA TGGCCGTGTG AACCGGTCTC TCGGACCGTA	
GTGAAGAGCT GCCCCATAT GACAGAGGAG TCTGCATCTC CCCAGAGGCT ATCGTCACAG	780
CACTTCTCGA CGGGCATATA CTGTCCTCTC AGACGTAGAG GGGTCTCCGA TAGCAGTGT	
TGGAAACAAGG AACAGATTCA ATGCCAGACT TCTCCATGGA TTCAAAACAAT GGAAATTGCG	840
ACCTTGTTC TTGTCTAAGT TACGGTCTGA AGAGGTACCT AAGTTGTTA CCTTTAACGC	
GAAGCGGCAG GGAGCACTGT AAATGCAAGC CCATGAAGGC AACCCAAAAG ACGTATCTCA	900
CTTCGCCGTC CCTCGTGTACA TTTACGTTCG GGTACTTCCG TTGGGTTTTC TGCACTAGAGT	
AGAATAATTAA CAATTATGTA ATCAGAGCAA AAGTGAAGAGA GGTGAAAGTG AAATGCCACG	960
TCTTATTAAAT GTTAATACAT TAGTCTCGTT TTCACTTCTC CCACCTTCAC TTTACGGTGC	
ACGCCAACAGC AATTGTGGAA GTAAAGGAGA TTCTCAAGTC TTCCCTAGTG AACATTCCCTA	1020
TGGCGTTGTGCG TTAACACCTT CATTCCCTCT AAGAGTTCAAG AAGGGATCAC TTGTAAGGAT	

AAGACACAGT GACACTGTAC ACCAACTCAG GCTGCTTGTG CCCCCAGCTT GTTGCCAATG TTCTGTGTCA CTGTGACATG TGGTTGAGTC CGACGAACAC GGGGGTCGAA CAACGGTTAC	1080
AGGAATACAT AATTATGGGC TATGAAGACA AAGAGCGTAC CAGGCTTCTA CTAGTGGAAG TCCTTATGTA TTAATACCCG ATACTTCTGT TTCTCGCATG GTCCGAAGAT GATCACCTTC	1140
GATCCTTGGC CGAAAAATGG AGAGATCGTC TTGCTAAGAA AGTCAGCGC TGGGATCAA CTAGGAACCG GCTTTTACCG TCTCTAGCAG AACGATTCTT TCAGTTCGCG ACCCTAGTTT	1200
AGCTTCGACG TCCCAGGAAA AGCAAAGACC CCGTGGCTCC AATTCCCAAC AAAAACAGCA TCGAAGCTGC AGGGTCCTT TCGTTTCTGG GGCACCGAGG TTAAGGGTTG TTTTGTGCGT	1260
ATTCCAGACA AGCGCGTAGT TAGACTAACG GAAAGGTGTA TGGAAACTCT ATGGACTTTG TAAGGTCTGT TCGCGCATCA ATCTGATTGC CTTTCCACAT ACCTTTGAGA TACCTGAAAC	1320
AAACTAAGAT TTGCATTGTT GGAAGAGCAA AAAAGAAATT GCACTACAGC ACGTTATATT TTTGATTCTA AACGTAACAA CCTTCTCGTT TTTCTTAA CGTGATGTG TGCAATATAA	1380
CTATTGTTA CTACAAGAAG CTGGTTAGT TGATTGAGT TCTCCTTCC TTCTTTTT GATAACAAAT GATGTTCTTC GACCAAATCA ACTAACATCA AGAGGAAAGG AAGAAAAAAA	1440
TTATAACTAT ATTTGCACGT GTTCCCAGGC AATTGTTTTA TTCAACTTCC AGTGACAGAG AATATTGATA TAAACGTGCA CAAGGGTCCG TTAACAAAAT AAGTTGAAGG TCACTGTCTC	1500
CAGTGACTGA ATGTCTCAGC CTAAAGAACG TCAATTCAATT TCTGATCAAC TAATGGTGAC GTCACTGACT TACAGAGTCG GATTCTTCG AGTTAAGTAA AGACTAGTTG ATTACCACTG	1560
AAAGTGTGTTGA TACTTGGGGA AAGTGAACTA ATTGCAATGG TAAATCAGAG AAAAGTTGAC TTCACAAACT ATGAACCCCT TTCACTTGAT TAACGTTACC ATTTAGTCTC TTTTCAACTG	1620
CAATGTTGCT TTTCCGTAG ATGAACAAGT GAGAGATCAC ATTTAAATGA TGATCAGCTT GTTACACGA AAAGGACATC TACTTGTCA CTCTCTAGTG TAAATTTACT ACTAGTGAAA	1680
CCATTTAATA CTTTCAGCAG TTTTAGTTAG ATGACATGTA GGATGCACCT AAATCTAAAT GGTAAATTAT GAAAGTCGTC AAAATCAATC TACTGTACAT CCTACGTGGA TTTAGATTAA	1740
ATTTTATCAT AAATGAAGAG CTGGTTAGA CTGTATGGTC ACTGTTGGGA AGGTAATGC TAAATAGTA TTTACTTCTC GACCAAATCT GACATACCG TGACAACCCCT TCCATTACG	1800
CTACTTTGTC AATTCTGTTT TAAAAATTGC CTAAATAAAT ATTAAGTCCT AAATAAAAAAA GATGAAACAG TTAAGACAAA ATTTTAACG GATTATTTA TAATTCAAGGA TTTATTTTTT	1860
AAAAAAAAAAA AAAAA TTTTTTTTTTT TTTTT	

Fig. 4. (Continuation page 2, SEQ ID NO:4).

MLLLFR A IPM	LLLGLMVIQT	DCEIAQYYID	EEEP P GT V IA	VLSQHSIFNT	TDIPATNFRL	60
MKQFNNSLIG	VRESDGQLSI	MERIDREQIC	RQSLHCNLAL	DVVSFSKGHF	KLLNVKVEVR	120
DINDHSPHFP	SEIMHVEVSE	SSSVGTRIPL	EIAIDEDVGS	NSIQNFQISN	NSHFSIDVLT	180
RADGVKYADL	VLMRELDREI	QPTYIMELLA	MDGGVPSLSG	TAVVNIRVLD	FNDNSPVFER	240
STIAV D LVED	APIGYLL L EL	HATDD D EGVN	GEIVYGFSTL	ASQEV R QLFK	INSRTGSVTL	300
EGQVDFETKQ	TYEF E VQAQD	LGP N PLTATC	KVT V HILDVN	DNTPAITITP	LTTVNAGVAY	360
IPETATKENF	IALISTTDRA	SGSNGQVRCT	LYGHEHFKLQ	QAYEDSYMIV	TTSTLDRENI	420
AAYSLTVVAE	DLGFP S LTKT	KYYTVKVSDE	NDNAPVFSKP	QYEASILENN	APGSYITT V I	480
ARDSDSDQNG	KVNYRLVDAK	VMGQSLTTFV	S I DADSGVLR	AVRS L DYEKL	KQLDFEIEAA	540
DNGIPQLSTR	VQLNLRIVDQ	NDNCPVITNP	LLNNNGS G EV L	LPISAPQNYL	V F QLKAEDSD	600
EGHNSQLFYT	ILRDPSRLFA	INKESGEVFL	KKQLNSDHSE	DLSTVVAVYD	LGRPSLSTNA	660
TVKFILTDSF	PSNVEVVILQ	PSAEEQHQID	MSIIFI A VLA	GGCALLLLAI	FFVACTCKKK	720
AGEFKQVPEQ	HGTCNEERLL	STPSPQSVSS	SLSQSESQL	SINTESENCS	VSSNQE Q HQQ	780
TGIKHSISVP	SYHTSGWHLD	NCAMSISGHS	HMGHISTKVQ	WAKEIVTSMT	VTLILVENQK	840
RRALSSQCRH	KPVLNTQMNQ	QGS D MPITIS	ATESTRVQKM	GTAHCNMKRA	IDCLTL	

Figure 5. Deduced amino acid sequence of the Xenopus PAPC (paraxial protocadherin) protein. It encodes a member of the cadherin family of transmembrane proteins that has dorsalizing activity when constructs are injected into Xenopus embryos. SEQ ID NO:5.

Figure 6. Nucleotide sequence of the full-length PAPC cDNA derived from the Xenopus organizer. The sense strand of the DNA is shown in the top line (in the 5' to 3' direction), and the bottom line shows the antisense strand (opposite orientation). SEQ ID NO:6.

GAATTCCCAG AGATGAACTC CTTGAGATTG TTTTAAATGA CTGCAGGTCT GGAAGGATTG	60
CTTAAGGGTC TCTACTTGAG GAACTCTAAC AAAATTTACT GACGTCCAGA CCTTCCTAAG	
ACATTGCCAC ACTGTTTCTA GGCAATGAAAA AACTGCAAGT TTCAACTTTG TTTTGGTGC	120
TGTAACGGTG TGACAAAGAT CCGTACTTT TTGACGTTCA AAGTTGAAAC AAAACCCACG	
AACTTGATT CTTCAAGATG CTGCTTCTCT TCAGAGCCAT TCCAATGCTG CTGTTGGAC	180
TTGAAACTAA GAAGTTCTAC GACGAAGAGA AGTCTCGGTA AGGTTACGAC GACAACCTG	
TGATGGTTT ACAAAACAGAC TGTGAAATTG CCCAGTACTA CATAGATGAA GAAGAACCCC	240
ACTACCAAA TGTTTGTCTG ACACTTTAAC GGGTCATGAT GTATCTACTT CTTCTGGGG	
CTGGCACTGT AATTGCAGTG TTGTACAAAC ACTCCATATT TAACACTACA GATATACCTG	300
GACCGTGACA TTAACGTCAC AACAGTGTG TGAGGTATAA ATTGTGATGT CTATATGGAC	
CAACCAATTG CCGTCTAATG AAGCAATTG ATAATTCCCT TATCGGAGTC CGTGAGAGTG	360
GTTGGTTAAA GGCAGATTAC TTCGTTAAAT TATAAGGGA ATAGCCTCAG GCACCTCAC	
ATGGGCAGCT GAGCATCATG GAGAGGATTG ACCGGGAGCA AATCTGCAGG CAGTCCTTC	420
TACCCGTCGA CTCGTAGTAC CTCTCTAAC TGGCCCTCGT TTAGACGTCC GTCAGGGAAAG	
ACTGCCACCT GGCTTTGGAT GTGGTCAGCT TTTCAAAGG ACACCTCAAG CTTCTGAACG	480
TGACGTTGGA CCGAAACCTA CACCAGTCGA AAAGGTTCC TGTGAAGTTC GAAGACTTGC	
TGAAAAGTGGG GGTGAGAGAC ATTAATGACC ATAGCCCTCA CTTTCCCAGT GAAATAATGC	540
ACTTCACCT CCACTCTCTG TAATTACTGG TATCAGGAGT GAAAGGGTCA CTTTATTACG	
ATGTGGAGGT GTCTGAAAGT TCCTCTGTGG GCACCAAGGAT TCCTTTAGAA ATTGCAATAG	600
TACACCTCCA CAGACTTTCA AGGAGACACC CGTGGTCCTA AGGAATCTT TAACGTTATC	
ATGAAGATGT TGGGTCCAAC TCCATCCAGA ACTTCAGAT CTCAAATAAT AGCCACTTCA	660
TACTTCTACA ACCCAGGTG AGGTAGGTCT TGAAAGTCTA GAGTTTATTA TCGGTGAAGT	
GCATTGATGT GCTAACCAAGA GCAGATGGGG TGAAATATGC AGATTTAGTC TTAATGAGAG	720
CGTAACTACA CGATTGGTCT CGTCTACCCC ACTTTATACG TCTAAATCAG AATTACTCTC	
AACTGGACAG GGAAATCCAG CCAACATACA TAATGGAGCT ACTAGCAATG GATGGGGGTG	780
TTGACCTGTC CCTTTAGGTC GGTTGTATGT ATTACCTCGA TGATCGTTAC CTACCCCCAC	
TACCATCACT ATCTGGTACT GCAGTGGTTA ACATCCGAGT CCTGGACTTT AATGATAACA	840
ATGGTAGTGA TAGACCATGA CGTCACCAAT TGTAGGCTCA GGACCTGAAA TTACTATTGT	
GCCCAGTGTG TGAGAGAAGC ACCATTGCTG TGGACCTAGT AGAGGATGCT CCTCTGGGAT	900
CGGGTCACAA ACTCTCTTCG TGGTAACGAC ACCTGGATCA TCTCCTACGA GGAGACCCCTA	
ACCTTTGTT GGAGTTACAT GCTACTGACG ATGATGAAGG AGTGAATGGA GAAATTGTTT	960
TGGAAAACAA CCTCAATGTA CGATGACTGC TACTACTTCC TCACTTACCT CTTAACAAA	
ATGGATTCAAG CACTTTGGCA TCTCAAGAGG TACGTCAAGCT ATTTAAAATT AACTCCAGAA	1020
TACCTAACGTC GTGAAACCGT AGAGTTCTCC ATGCAGTCGA TAAATTTAA TTGAGGTCTT	

CTGGCAGTGT TACTCTTGAA GGCCAAGTTG ATTTGAGAC CAAGCAGACT TACGAATTTG GACCGTCACA ATGAGAACCTT CCGGTTCAAC TAAAACCTTG GTTCGTCGAA ATGCTTAAAC	1080
AGGTACAAGC CCAAGATTTG GGCCCCAACCC CACTGACTGC TACTTGTAAA GTAACTGTTC TCCATGTTCG GGTTCTAAAC CCGGGGTTGG GTGACTGACG ATGAACATT CATTGACAAG	1140
ATATACTTGA TGTAATGAT AATACCCCCAG CCATCACTAT TACCCCTCTG ACTACTGTAA TATATGAAC TACATTTACTA TTATGGGTC GGTAGTGATA ATGGGGAGAC TGATGACATT	1200
ATGCAGGAGT TGCCTATATT CCAGAAACAG CCACAAAGGA GAACTTTATA GCTCTGATCA TACGTCCCTCA ACGGATATAAA GGTCTTGTC GGTGTTCCCT CTTGAAATAT CGAGACTAGT	1260
GCACACTGAA CAGAGCCTCT GGATCTAATG GACAAGTTCG CTGTAACCTT TATGGACATG CGTGATGACT GTCTCGGAGA CCTAGATTAC CTGTTCAAGC GACATGAGAA ATACCTGTAC	1320
AGCACTTAA ACTACAGCAA GCTTATGAGG ACAGTTACAT GATAGTTACC ACCTCTACTT TCGTGAAATT TGATGTCGTT CGAATACTCC TGTCAATGTA CTATCAATGG TGGAGATGAA	1380
TAGACAGGGA AAACATAGCA GCGTACTCTT TGACAGTAGT TGCAGAAGAC CTTGGCTTCC ATCTGTCCTT TTTGTATCGT CGCATGAGAA ACTGTCATCA ACGTCTTCTG GAACCGAAGG	1440
CCTCATTGAA GACCAAAAAG TACTACACAG TCAAGGTTAG TGATGAGAAAT GACAATGCAC GGAGTAACCTT CTGGTTTTTC ATGATGTCG AGTTCCAATC ACTACTCTTA CTGTTACGTG	1500
CTGTATTTTC TAAACCCCCAG TATGAAGCTT CTATTCTGGA AAATAATGCT CCAGGCTCTT GACATAAAAG ATTTGGGTC ATACTTCGAA GATAAGACCT TTTATTACGA GGTCCGAGAA	1560
ATATAACTAC AGTGATAGGCC AGAGACTCTG ATAGTGATCA AAATGGCAA GTAAATTACA TATATTGATG TCACTATCGG TCTCTGAGAC TATCACTAGT TTTACCGTTT CATTAAATGT	1620
GACTTGTGGA TGCAAAAGTG ATGGGCCAGT CACTAACAAAC ATTTGTTCT CTTGATGCGG CTGAACACCT ACGTTTCAC TACCCGGTCA GTGATTGTTG TAAACAAAGA GAACTACGCC	1680
ACTCTGGAGT ATTGAGAGCT GTTAGGTCTT TAGACTATGA AAAACTTAA CAACTGGATT TGAGACCTCA TAACTCTCGA CAATCCAGAA ATCTGATACT TTTTGAATT GTGACCTAA	1740
TTGAAATTGA AGCTGCAGAC AATGGGATCC CTCAACTCTC CACTCGCGTT CAACTAAATC AACTTTAACT TCGACGTCTG TTACCCCTAGG GAGTTGAGAG GTGAGCGCAA GTTGATTTAG	1800
TCAGAAATGT TGATCAAAAT GATAATTGCC CTGTGATAAC TAATCCTCTT CTTAATAATG AGTCTTATCA ACTAGTTTTA CTATTAACGG GACACTATTG ATTAGGAGAA GAATTATTAC	1860
GCTCGGGTGA AGTTCTGCTT CCCATCAGCG CTCCCTAAAA CTATTTAGTT TTCCAGCTCA CGAGCCCCACT TCAAGACGAA GGGTAGTCGC GAGGAGTTT GATAAATCAA AAGGTCGAGT	1920
AAGCCGAGGA TTCAGATGAA GGGCACAACT CCCAGCTGTT CTATACCATCA CTGAGAGATC TTCGGCTCCT AAGTCTACTT CCCGTGTTGA GGGTCGACAA GATATGGTAT GACTCTCTAG	1980
CAAGCGAGATT GTTTGCCATT AACAAAGAAA GTGGTGAAAGT GTTCCTGAAA AAACAATTAA GTTCGTCTAA CAAACGGTAA TTGTTCTTT CACCACTTCA CAAGGACTTT TTTGTTAATT	2040
ACTCTGACCA TTCAGAGGAC TTGAGCATAG TAGTTGCGAT GTATGACTTG GGAAGACCTT TGAGACTGGT AAGTCTCCTG AACTCGTATC ATCAACGTCA CATACTGAAC CCTTCTGGAA	2100
CATTATCCAC CAATGCTACA GTTAAATTCA TCCTCACCGA CTCTTTCCCT TCTAACGTTG GTAATAGGTG GTTACGATGT CAATTAAAGT AGGAGTGGCT GAGAAAAGGA AGATTGCAAC	2160

Fig. 6. (Continuation page 2, SEQ ID NO:6).

AGTCGTTAT TTTGCAACCA TCTGCAGAAG AGCAGCACCA GATCGATATG TCCATTATAT	2220
TTCAGCAATA AAACGTTGGT AGACGTCTTC TCGTCGTGGT CTAGCTATAC AGGTAATATA	
TCATTCGAGT GCTGGCTGGT GGTTGTGCTT TGCTACTTTT GGCCATCTTT TTTGTGGCCT	2280
AGTAACGTCA CGACCGACCA CCAACACGAA ACGATGAAAA CCGGTAGAAA AAACACCGGA	
GTACTTGTAA AAAGAAAGCT GGTGAATTAA AGCAGGTACC TGAACAACAC GGAACATGCA	2340
CATGAACATT TTTCTTCGA CCACTTAAAT TCGTCCATGG ACTTGTGTTG CCTTGTACGT	
ATGAAGAACG CCTGTTAACGC ACCCCATCTC CCCAGTCGGT CTCTTCTTCT TTGTCAGT	2400
TACTTCTTGC GGACAATTG TGTTGAGAGG GGGTCAGCCA GAGAAGAAGA AACAGAGTCA	
CTGAGTCATG CCAACTCTCC ATCAAACTG AATCTGAGAA TTGCAGCGTG TCCTCTAAC	2460
GACTCAGTAC GGTTGAGAGG TAGTTATGAC TTAGACTCTT AACGTCGCAC AGGAGATTGG	
AAGAGCAGCA TCAGCAAACA GGCATAAAGC ACTCCATCTC TGTACCATCT TATCACACAT	2520
TTCTCGTCGT AGTCGTTTGT CCGTATTTCG TGAGGTAGAG ACATGGTAGA ATAGTGTGTA	
CTGGTTGGCA CCTGGACAAT TGTGCAATGA GCATAAGTGG ACATTCTCAC ATGGGGCACA	2580
GACCAACCGT GGACCTGTAA ACACGTTACT CGTATTCCACC TGTAAGAGTG TACCCCGTGT	
TTAGTACAAA GGTACAGTGG GCAAAGGAGA TAGTGACTTC AATGACAGTG ACTCTGATAC	2640
AATCATGTTT CCATGTCACC CGTTTCTCT ATCACTGAAG TTACTGTCAC TGAGACTATG	
TAGTGGAGAA TCAGAAAAGA AGAGCATTGA GCAGCCAATG CAGGCACAAG CCAGTGCTCA	2700
ATCACCTCTT AGTCTTTCT TCTCGTAACT CGTCGGTTAC GTCCGTGTTG GGTACCGAGT	
ATACACAGAT GAATCAGCAG GGTTCCGACA TGCCGATAAC TATTCAGCC ACCGAATCAA	2760
TATGTGTCTA CTTAGTCGTC CCAAGGCTGT ACGGCTATTG ATAAAGTCGG TGGCTTAGTT	
CAAGGGTCCA GAAAATGGGA ACTGCACATT GCAATATGAA AAGGGCTATA GACTGTCTTA	2820
GTTCCCAGGT CTTTACCCCT TGACGTGTAA CGTTACTATT TTCCCGATAT CTGACAGAAT	
CTCTGTAGCT CCTGTATATT ACAATACCTA CCATGCAAGA ATGCCCTAACCC TGCACATACC	2880
GAGACATCGA GGACATATAA TGTTATGGAT GGTACGTTCT TACGGATTGG ACGTGTATGG	
GAACCATACC CTTAGAGACC CTTATTACCA TATCAATAAT CCTGTTGCTA ATCGGATGCA	2940
CTTGGTATGG GAATCTCTGG GAATAATGGT ATAGTTATTA GGACAAACGAT TAGCCTACGT	
GGCGGAATAT GAAAGAGATT TAGTCAACAG AAGTGCACCG TTATCTCCGC AGAGATCGTC	3000
CCGCCTTATA CTTTCTCTAA ATCAGTTGTC TTCACGTTGC AATAGAGGCG TCTCTAGCAG	
TAGCAGATAC CAAGAATTCA ATTACAGTCC GCAGATATCA AGACAGCTTC ATCCCTCAGA	3060
ATCGTCTATG GTTCTTAAGT TAATGTCAGG CGTCTATAGT TCTGTCGAAG TAGGAAGTCT	
AATTGCTACA ACCTTTAAT CATTAGGCAT GCAAGTGAGA ATGCACAAAG GCAAGTGCTT	3120
TTAACGATGT TGGAAAATTA GTAATCCGTA CGTTCACTCT TACGTGTTTC CGTTCACGAA	
TAGCATGAAA GCTAAATATA TGGAGTCTCC CCTTTCCCTC TGATGGATGG GGGGAGACAC	3180
ATCGTACTTT CGATTTATAT ACCTCAGAGG GGAAAGGGAG ACTACCTACC CCCCTCTGTG	
AGGACAGTGC ATAAATATAC AGCTGTTTC TATTTGCATT TCACGGAGA ATTTTTGTT	3240
TCTCTGTCACG TATTTATATG TCGACGAAAG ATAAACGTAAG AGTGAACCCCT TAAAAAACAA	
TTTTTACAT ATTTATTTT CCTGAATTGA ATGTGACATT GTCCTGTCAC CTAACCTAGCA	3300
AAAAAATGTA TAAATAAAAA GGACTTAAC TACACTGAA CAGGACAGTG GATTGATCGT	

Fig. 6. (Continuation page 3, SEQ ID NO:6).

ATTAATCCA CAGACCTACA GTCAAATATT TGAGGGCCCC TGAAACAGCA CATCAGTCAG TAATTTAGGT GTCTGGATGT CAGTTATAA ACTCCCGGGG ACTTTGTCGT GTAGTCAGTC	3360
GACCTAAAGT GGCTTTTTA CTTTAGCAG CTCTGGTC TGCCCTCTGT GTTAATCAGC CTGGATTCA CGGGAAAAT GAAAATCGTC GAGGACCCAG ACGGGAGACA CAATTAGTCG	3420
CCCTGGTCAA GTCTGAGTA GGATCATGGC GTTTTATAT GCATCTCACC TACTTTGGAC GGGACCAAGT CAGGACTCAT CCTAGTACCG CAAAAATATA CGTAGAGTGG ATGAAACCTG	3480
GTGATTTACA CATAATAGGA AACGCTGGT TTCAGTGAAG TCTGTGTTGT ATATATTCTG CACTAAATGT GTATTATCCT TTGCGAACCA AAGTCACTTC AGACACAACA TATATAAGAC	3540
TTATATACAC GCATTTGTG TTTGTGTATA TATTCAAGT CCATTCAGAT ATGTGTATAT AATATATGTG CGTAAACAC AAACACATAT ATAAAGTTCA GGTAAGTCTA TACACATATA	3600
AGTGCAGACC TTGTAAATTA AATATTCTGA TACTTTTCC TCAATAAATA TTTAAAT TCACGTCTGG AACATTTAAT TTATAAGACT ATGAAAAAGG AGTTATTAT AAATTTA	

Fig. 6. (Continuation page 4, SEQ ID NO:6).

MVCCGPGRML LGWAGLLVLA ALCLLQVPGQ QAAACEPVRI PLCKSLPWNM TKMPNHLHHS	60
TQANAILAME QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE	120
PILIKYRHSW PESLACDELP VYDRGVCISP EAIVTADGAD FPMDSSTGHC RGASSERCKC	180
KPVRATQKTY FRNNNYNYVIR AKVKEVKMKC HDVTAVVEVK EILKASLVNI PRDTVNLYTT	240
SGCLCPPLTV NEEYVIMGYE DEERSRLLLVEGSIAEKWKD RLGKKVKRWD MKLRHLGLGK	300
TDASDSTQNZ KSGRNSNPRP ARS.	

Figure 7. Deduced amino acid sequence of mouse FRZB-1 protein. SEQ ID NO:7.

Figure 8. Nucleotide sequence of the full-length mouse FRZB-1 cDNA. SEQ ID NO:8.

AAGCCTGGGA CCATGGTCTG CTGCGGCCCG GGACGGATGC TGCTAGGATG GGCCGGGTTG TTCGGACCT GGTACCAGAC GACGCCGGC CCTGCCTACG ACGATCCTAC CGGGCCCAAC	60
CTAGTCCTGG CTGCTCTCTG CCTGCTCCAG GTGCCCGGAG CTCAGGCTGC AGCCTGTGAG GATCAGGACC GACGAGAGAC GGACGAGGTC CACGGGCCTC GAGTCCGACG TCGGACACTC	120
CCTGTCCGCA TCCCGCTGTG CAAGTCCCTT CCCTGGAACA TGACCAAGAT GCCCAACCAC GGACAGGCCT AGGGCGACAC GTTCAGGGAA GGGACCTTGT ACTGGTTCTA CGGGTTGGTG	180
CTGCACCAACA GCACCCAGGC TAACGCCATC CTGGCCATGG AACAGTCGA AGGGCTGCTG GACGTGGTGT CGTGGGTCCG ATTGCGGTAG GACCGGTACC TTGTCAAGCT TCCCAGCGAC	240
GGCACCCACT GCAGCCCGGA TCTTCTCTTC TTCCCTGTG CAATGTACGC ACCCATTG CCGTGGGTGA CGTCGGGCCT AGAAGAGAAG AAGGAGACAC GTTACATGCG TGGGTAAACG	300
ACCATCGACT TCCAGCACGA GCCCATCAAG CCCTGCAAGT CTGTGTGTGA GCGCGCCCGA TGGTAGCTGA AGGTCGTGCT CGGGTAGTTC GGGACGTTCA GACACACACT CGCGCGGGCT	360
CAGGGCTGCG AGCCCATTCT CATCAAGTAC CGCCACTCGT GGCCGGAAAG CTTGGCCTGC GTCCCGACGC TCGGGTAAGA GTAGTTCATG GCGGTGAGCA CCGGCCTTTC GAACCGGACG	420
GACGAGCTGC CGGTGTACGA CCGCGCGGTG TGCATCTCTC CTGAGGCCAT CGTCACCGCG CTGCTCGACG GCCACATGCT GGCGCCGCAC ACGTAGAGAG GACTCCGGTA GCAGTGGCGC	480
GACGGAGCGG ATTTTCCTAT GGATTCAAGT ACTGGACACT GCAGAGGGGC AAGCAGCGAA CTGCCTCGCC TAAAAGGATA CCTAAGTTCA TGACCTGTGA CGTCTCCCCG TTCGTGCGTT	540
CGTTGCAAAT GTAAGCCTGT CAGAGCTACA CAGAAGACCT ATTTCCGGAA CAATTACAAC GCAACGTTA CATTGGACA GTCTCGATGT GTCTTCTGGAA TAAAGGCCTT GTTAATGTTG	600
TATGTCATCC GGGCTAAAGT TAAAGAGGTA AAGATGAAAT GTCATGATGT GACCGCCGTT ATACAGTAGG CCCGATTCA ATTTCTCCAT TTCTACTTTA CAGTACTACA CTGGCGCAA	660
GTGGAAGTGA AGGAAATTCT AAAGGCATCA CTGGTAAACA TTCCAAGGGAA CACCGTCAAT CACCTCACT TCCTTTAAGA TTTCCGTAGT GACCATTGT AAGGTTCCCT GTGGCAGTTA	720
CTTTATACCA CCTCTGGCTG CCTCTGTCTT CCACTTACTG TCAATGAGGA ATATGTCATC GAAATATGGT GGAGACCGAC GGAGACAGGA GGTGAATGAC AGTTACTCCT TATACAGTAG	780
ATGGGCTATG AAGACGAGGA ACGTTCCAGG TTACTCTTGG TAGAAGGCTC TATAGCTGAG TACCCGATAC TTCTGCTCCT TGCAAGGTCC AATGAGAACCC ATCTTCCGAG ATATCGACTC	840
AAGTGGAAAGG ATCGGCTTGG TAAGAAAGTC AAGCGCTGGG ATATGAAACT CCGACACCTT TTCACCTTCC TAGCCGAACC ATTCTTTCAG TTCGCGACCC TATACTTTGA GGCTGTGGAA	900
GGACTGGGTA AAACTGATGC TAGCGATTCC ACTCAGAACATC AGAAGTCTGG CAGGAACCT CCTGACCCAT TTTGACTACG ATCGCTAAGG TGAGTCTTAG TCTTCAGACC GTCCTTGAGA	960

AATCCCCGGC CAGCACGCAG CCAAATCCTG AAATGTAAAA GGCCACACCC ACGGACTCCC TTAGGGGCCG GTCGTGCGTC GATTTAGGAC TTTACATTTT CCGGTGTGGG TGCCTGAGGG	1020
TTCTAAGACT GGCGCTGGTG GACTAACAAA GGAAAACCGC ACAGTTGTGC TCGTGACCGA AAGATTCTGA CGCGACAC CTGATTGTTT CCTTTGGCG TGTCAACACG AGCACTGGCT	1080
TTGTTTACCG CAGACACCGC GTGGCTACCG AAGTTACTTC CGGTCCCTT TCTCCTGCTT AACAAATGGC GTCTGTGGCG CACCGATGGC TTCAATGAAG GCCAGGGAA AGAGGACGAA	1140
CTTAATGGCG TGGGTTAGA TCCTTTAATA TGTTATATAT TCTGTTTCAT CAATCACGTG GAATTACCGC ACCCCAAATCT AGGAAATTAT ACAATATATA AGACAAAGTA GTTAGTGCAC	1200
GGGACTGTTC TTTTGCAACC AGAATAGTAA ATTAAATATG TTGATGCTAA GGTTCTGTA CCCTGACAAG AAAACGTTGG TCTTATCATT TAATTTATAC AACTACGATT CCAAAGACAT	1260
CTGGACTCCC TGGGTTTAAT TTGGTGTCT GTACCCGTAT TGAGAATGCA ATGTTTCATG GACCTGAGGG ACCCAAATTA AACACACAAGA CATGGGACTA ACTCTTACGT TACAAAGTAC	1320
TAAAGAGAGA ATCCTGGTCA TATCTCAAGA ACTAGATATT GCTGTAAGAC AGCCTCTGCT ATTCTCTCT TAGGACCAGT ATAGAGTTCT TGATCTATAA CGACATTCTG TCGGAGACGA	1380
GCTGCGCTTA TAGTCTTGTG TTTGTATGCC TTTGTCCATT TCCCTCATGC TGTGAAAGTT CGACGCGAAT ATCAGAACAC AAACATACGG AAACAGGTAA AGGGAGTACG ACACTTCAA	1440
ATACATGTTT ATAAAGGTAG AACGGCATT TGAAATCAGA CACTGCACAA GCAGAGTAGC TATGTACAAA TATTTCCATC TTGCCGTAAA ACTTTAGTCT GTGACGTGTT CGTCTCATCG	1500
CCAACACCAG GAAGCATTAA TGAGGAAACG CCACACAGCA TGACTTATT TCAAGATTGG GGTTGTGGTC CTTCGTAAAT ACTCCCTTGC GGTGTGCGT ACTGAATAAA AGTTCTAAC	1560
CAGGCAGCAA AATAAAATAGT GTTGGGAGCC AAGAAAAGAA TATTTTGCCCT GGTTAAGGGG GTCCGTCGTT TTATTTATCA CAACCCCTCGG TTCTTTCTT ATAAAACGGA CCAATTCCCC	1620
CACACTGGAA TCAGTAGGCC TTGAGCCATT AACAGCAGTG TTCTTCTGGC AAGTTTTGA GTGTGACCTT AGTCATCGGG AACTCGGTAA TTGTCGTCAC AAGAAGACCG TTCAAAAAC	1680
TTTGTTCATA AATGTATTCA CGAGCATTAG AGATGAACCTT ATAACTAGAC ATCTGTTGTT AAACAAGTAT TTACATAAGT GCTCGTAATC TCTACTTGAA TATTGATCTG TAGACAACAA	1740
ATCTCTATAG CTCTGCTTCC TTCTAAATCA AACCCATTGT TGGATGCTCC CTCTCCATT TAGAGATATC GAGACGAAGG AAGATTTAGT TTGGGTAAACA ACCTACGAGG GAGAGGTAAG	1800

ATAAATAAAT TTGGCTTGCT GTATTGGCCA GGAAAAGAAA GTATTAAAGT ATGCATGCAT 1860
TATTTATTTA AACCGAACGA CATAACCGGT CCTTTCTTT CATAATTCA TACGTACGTA

GTGCACCAGG GTGTTATTTA ACAGAGGTAT GTAACTCTAT AAAAGACTAT AATTACAGG 1920
CACGTGGTCC CACAATAAAT TGTCTCCATA CATTGAGATA TTTTCTGATA TTAAATGTCC

ACACGGAAAT GTGCACATTT GTTACTTTT TTTCTCCCTT TTGCTTGAGG CTTGTGATTT 1980
TGTGCCTTTA CACGTGTAAA CAAATGAAAA AAAGAAGGAA AACGAAACCC GAACACTAAA

TGGTTTTTGG TGTGTTATG TCTGTATTTT GGGGGGTGGG TAGGTTTAAG CCATTGCACA 2040
ACCAAAAACC ACACAAATAC AGACATAAAA CCCCCCACCC ATCCAAATTG GGTAACGTGT

TTCAAGTTGA ACTAGATTAG AGTAGACTAG GCTCATTGGC CTAGACATTA TGATTGAAT 2100
AAGTTCAACT TGATCTAACATC TCATCTGATC CGAGTAACCG GATCTGTAAT ACTAAACTTA

TTGTGTTGTT TAATGCTCCA TCAAGATGTC TAATAAAAGG AATATGGTG TCAACAGAGA 2160
AACACAAACAA ATTACGAGGT AGTTCTACAG ATTATTTCC TTATACCAAC AGTTGTCTCT

CGACAACAAC AACAAA
GCTGTTGTTG TTGTTT

MVCGSPGGML LLRAGLLALA ALCLLRVPGA RAAACEPVRI PLCKSLPWNM TKMPNHLHHS	60
TQANAILAIE QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE	120
PILIKYRHSW PENLACEELP VYDRGVCISP EAIVTADGAD FPMDSNGNC RGASSERCKC	180
KPIRATQKTY FRNNNYNYVIR AKVKEIKTKC HDVTAVVEVK EILKSSLVNI PRDTVNLYTS	240
SGCLCPPLNV NEEYIIMGYE DEERSRLLL V EGSIAEKWKD RLGKKVKRWD MKLRHLGLSK	300
SDSSNSDSTQ SQKSGRNSNP RQARN.	

Figure 9. Deduced amino acid sequence of human FRZB-1 protein. SEQ ID NO:9.

Figure 10. Nucleotide sequence of the full-length human FRZB-1 cDNA. SEQ ID NO:10.

This sequence was assembled from public ESTs from the Genbank database (accession numbers: H18848, R63748, W38677, W44760, H38379 and N71244).

GGCGGAGCGG	GCCTTTGGC	GTCCACTGCG	CGGCTGCACC	CTGCCCCATC	TGCCGGGATC	60
CCGCCTCGCC	CGGAAAACCG	CAGGTGACGC	GCCGACGTGG	GACGGGGTAG	ACGGCCCTAG	
ATGGTCTGCG	GCAGCCGGG	AGGGATGCTG	CTGCTGCGGG	CCGGGCTGCT	TGCCCTGGCT	120
TACCAAGACGC	CGTCGGGCC	TCCCTACGAC	GACGACGCC	GGCCCGACGA	ACGGGACCGA	
GCTCTCTGCC	TGCTCCGGT	GCCCGGGGCT	CGGGCTGCAG	CCTGTGAGCC	CGTCCGCATC	180
CGAGAGACGG	ACGAGGCCA	CGGGCCCCGA	GCCCGACGTC	GGACACTCGG	GCAGGGTAG	
CCCCTGTGCA	AGTCCCTGCC	CTGGAACATG	ACTAAGATGC	CCAACCACCT	GCACCCACAGC	240
GGGGACACGT	TCAGGGACGG	GACCTTGTAC	TGATTCTACG	GGTTGGTGGA	CGTGGTGTG	
ACTCAGGCCA	ACGCCATCCT	GGCCATCGAG	CAGTCGAAG	GTCTGCTGGG	CACCCACTGC	300
TGAGTCCGGT	TGCGGTAGGA	CCGGTAGCTC	GTCAAGCTTC	CAGACGACCC	GTGGGTGACG	
AGCCCCGATC	TGCTCTCTT	CCTCTGTGCC	ATGTACGCGC	CCATCTGCAC	CATTGACTTC	360
TCGGGGCTAG	ACGAGAAAGAA	GGAGACACGG	TACATGCGCG	GGTAGACGTG	GTAACTGAAG	
CAGCACGAGC	CCATCAAGCC	CTGTAAGTCT	GTGTGCGAGC	GGGCCCGGCA	GGGCTGTGAG	420
GTCGTGCTCG	GGTAGTCGG	GACATTCAAGA	CACACGCTCG	CCCAGGGCCGT	CCCGACACTC	
CCCATACTCA	TCAAGTACCG	CCACTCGTGG	CCGGAGAACCC	TGGCCTGCCA	GGAGCTGCCA	480
GGGTATGAGT	AGTTCATGGC	GGTGAGCACC	GGCCTCTTGG	ACCGGACGCT	CCTCGACGGT	
GTGTACGACA	GGGGCGTGTG	CATCTCTCCC	GAGGCCATCG	TTACTGCGGA	CGGAGCTGAT	540
CACATGCTGT	CCCCGCACAC	GTAGAGAGGG	CTCCGGTAGC	AATGACGCCT	GCCTCGACTA	
TTTCCTATGG	ATTCTAGTAA	CGGAAACTGT	AGAGGGCAA	GCAGTGAACG	CTGTAAATGT	600
AAAGGATAACC	TAAGATCATT	GCCTTTGACA	TCTCCCCGTT	CGTCACTTGC	GACATTACCA	
AAGCCTATTA	GAGCTACACA	GAAGACCTAT	TTCCGGAACA	ATTACAACCA	TGTCATTGG	660
TTCCGATAAT	CTCGATGTGT	CTTCTGGATA	AAGGCCCTGT	TAATGTTGAT	ACAGTAAGCC	
GCTAAAGTTA	AAGAGATAAA	GACTAAGTGC	CATGATGTGA	CTGCAGTAGT	GGAGGTGAAG	720
CGATTTCAAT	TTCTCTATTT	CTGATTCAAG	GTACTACACT	GACGTCACTCA	CCTCCACTTC	
GAGATTCTAA	AGTCCTCTCT	GGTAAACATT	CCACGGGACA	CTGTCAACCT	CTATACCAGC	780
CTCTAAGATT	TCAGGAGAGA	CCATTGTAA	GGTGCCTGT	GACAGTTGGA	GATATGGTCG	
TCTGGCTGCC	TCTGCCCTCC	ACTTAATGTT	AATGAGGAAT	ATATCATCAT	GGGCTATGAA	840
AGACCGACGG	AGACGGGAGG	TGAATTACAA	TTACTCCTTA	TATAGTAGTA	CCCGATACTT	

GATGAGGAAC	GTTCCAGATT	ACTCTGGTG	GAAGGCTCTA	TAGCTGAGAA	GTGGAAGGAT	900
CTACTCCTTG	CAAGGTCTAA	TGAGAACAC	CTTCCGAGAT	ATCGACTCTT	CACCTTCCTA	
CGACTCGGTA	AAAAAGTTAA	GCGCTGGGAT	ATGAAGCTTC	GTCATCTTGG	ACTCAGTAAA	960
GCTGAGCCAT	TTTTTCAATT	CGCGACCCTA	TACTTCGAAG	CAGTAGAACCC	TGAGTCATTT	
AGTGATTCTA	GCAATAGTGA	TTCCACTCAG	AGTCAGAAGT	CTGGCAGGAA	CTCGAACCCC	1020
TCACTAAGAT	CGTTATCACT	AAGGTGAGTC	TCAGTCTTCA	GACCCTCCTT	GAGCTTGGGG	
CGGCAAGCAC	GCAACTAAAT	CCCGAAATAC	AAAAAGTAAC	ACAGTGGACT	TCCTATTAAG	1080
GCCGTTCGTG	CGTTGATTAA	GGGCTTTATG	TTTTTCATTG	TGTCACCTGA	AGGATAATTC	
ACTTACTTGC	ATTGCTGGAC	TAGCAAAGGA	AAATTGCACT	ATTCACATC	ATATTCTATT	1140
TGAATGAACG	TAACGACCTG	ATCGTTTCCT	TTTAACGTGA	TAACGTGTAG	TATAAGATAA	
GTTTACTATA	AAAATCATGT	GATAACTGAT	TATTACTTCT	GTTCCTCTTT	TGGTTTCTGC	1200
CAAATGATAT	TTTTAGTACA	CTATTGACTA	ATAATGAAGA	CAAAGAGAAA	ACCAAAGACG	
TTCTCTCTTC	TCTCAACCCC	TTTGTAAATGG	TTTGGGGGCA	GAATCTTAAG	TATATTGTGA	1260
AAGAGAGAAG	AGAGTTGGGG	AAACATTACC	AAACCCCGT	CTGAGAACATC	ATATAACACT	
GTTTTCTATT	TCACTAATCA	TGAGAAAAAC	TGTTCTTTG	CAATAATAAT	AAATTAAACA	1320
CAAAAGATAA	AGTGATTAGT	ACTCTTTTG	ACAAGAAAAC	GTTATTATTA	TTTAATTGTG	
TGCTGTTACC	AGAGCCTCTT	TGCTGAGTCT	CCAGATGTTA	ATTTACTTTC	TGCACCCCAA	1380
ACGACAATGG	TCTCGGAGAA	ACGACTCAGA	GGTCTACAAT	AAATGAAAG	ACGTGGGTT	
TTGGGAATGC	AATATTGGAT	AAAAAGAGAG	GTTCCTGGTA	TTCACAGAAA	GCTAGATATG	1440
AACCCCTTACG	TTATAACCTA	CTTTTCTCTC	CAAAGACCAT	AAAGTGTCTTT	CGATCTATAC	
CCTTAAAACA	TACTCTGCCG	ATCTAATTAC	AGCCTTATTT	TTGTATGCCT	TTTGGCATT	1500
GGAAATTGTG	ATGAGACGGC	TAGATTAATG	TCGGAATAAA	AAACATACGGA	AAACCCGTAA	
CTCCTCATGC	TTAGAAAGTT	CCAAATGTTT	ATAAAGGTAA	AAATGGCAGTT	TGAAGTCAAA	1560
GAGGAGTACG	AACTTTCAA	GGTTTACAAA	TATTTCCATT	TTACCGTCAA	ACTTCAGTTT	
TGTCACATAG	GCAAAGCAAT	CAAGCACCAAG	GAAGTGTGTTA	TGAGGAAACA	ACACCCAAGA	1620
ACAGTGTATC	CGTTTCGTTA	GTTCGTGGTC	CTTCACAAAT	ACTCCTTTGT	TGTGGGTTCT	
TGAATTATTT	TTGAGACTGT	CAGGAAGTAA	AATAAATAGG	AGCTTAAGAA	AGAACATTAA	1680
ACTTAATAAA	AACTCTGACA	GTCCTTCATT	TTATTTATCC	TCGAATTCTT	TCTTGTAAAA	
GCCTGATTGA	GAAGCACAAAC	TGAAACCAGT	AGCCGCTGGG	GTGTTAATGG	TAGCATTCTT	1740
CGGACTAACT	CTTCGTGTTG	ACTTTGGTCA	TCGGCGACCC	CACAATTACC	ATCGTAAGAA	
CTTTTGGCAA	TACATTGAT	TTGTTCATGA	ATATATTAAT	CAGCATTAGA	GAAATGAATT	1800
GAAAACCGTT	ATGTAACACTA	AACAAGTACT	TATATAATTA	GTCGTAATCT	CTTTACTTAA	
ATAAACTAGAC	ATCTGCTGTT	ATCACCATAG	TTTGTTTAA	TTTGCTTCCT	TTTAAATAAA	1860
TATTGATCTG	TAGACGACAA	TAGTGGTATC	AAAACAAATT	AAACGAAGGA	AAATTATTT	
CCCATTGGTG	AAAGTCAAAA	AAAAAAAAAA	AAA			
GGGTAAACCAC	TTTCAGTTTT	TTTTTTTTTT	TTT			